

Wang

1635

11/5/98

CRF Errors Corrected by the STIC System Branch

CRF Processing Date: \_\_\_\_\_  
Edited by: \_\_\_\_\_  
Verified by: \_\_\_\_\_ (STIC staff)

Serial Number: 08/978,636

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: **ENTERED**
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;  
☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/978,636DATE: 11/05/98  
TIME: 14:41:32

INPUT SET: S29661.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

## SEQUENCE LISTING

1  
2  
3 (1) General Information  
4  
5 (i) APPLICANT: ELAZAR RABBANI  
6 JANNIS G. STAVRIANOPOULOS  
7 JAMES J. DONEGAN  
8 DAKAI LIU  
9 NORMAN E. KELKER  
10 DEAN L. ENGELHARDT  
11  
12 (ii) TITLE OF INVENTION: NOVEL PROPERTY EFFECTING  
13 AND/OR PROPERTY EXHIBITING COMPOSITIONS FOR  
14 THERAPEUTIC AND DIAGNOSTIC USE  
15  
16 (iii) NUMBER OF SEQUENCES: 42  
17  
18 (iv) CORRESPONDENCE ADDRESS:  
19 (A) ADDRESSEE: ENZO THERAPEUTICS, INC.  
20 (B) STREET: C/O ENZO BIOCHEM, INC.  
21 527 MADISON AVENUE,  
22 9TH FLOOR  
23 (C) CITY: NEW YORK  
24 (D) STATE: NY  
25 (E) COUNTRY: USA  
26 (F) ZIP: 10022  
27  
28 (v) COMPUTER READABLE FORM:  
29 (A) MEDIUM TYPE: 3.5" Micro Floppy Disk. 1.44 KB  
30 STORAGE  
31 (B) COMPUTER: IBM PC/XT/AT, IBM PS/2 OR COMPATIBLES  
32 (C) OPERATING SYSTEM: PC-DOS  
33 (D) SOFTWARE: MICROSOFT WORD \_\_\_\_ - ASCII TEXT (DOS)  
34  
35 (vi) CURRENT APPLICATION DATA:  
36 (A) APPLICATION NUMBER: US 08/978,636  
37 (B) FILING DATE: 25-NOVEMBER-1997  
--> 38 (C) CLASSIFICATION: Not Yet Known  
39  
40 (viii) ATTORNEY/AGENT INFORMATION:  
41 (A) NAME: FEDUS, RONALD C.  
42 (B) REGISTRATION NUMBER: 32,567  
43 (C) REFERENCE/DOCKET NUMBER: ENZ-53(D3)  
44  
45 (ix) TELECOMUNICATION INFORMATION:

INPUT SET: S29661.raw

46 (A) TELEPHONE: (212) 583-0100  
47 (B) TELEFAX: (212) 583-0150  
48

## ERRORED SEQUENCES FOLLOW:

814 (2) INFORMATION FOR SEQ ID NO:42:  
815  
816 (i) SEQUENCE CHARACTERISTICS:  
--> 817 (A) LENGTH:67 base pairs  
818 (B) TYPE:nucleic acid  
819 (C) STRANDEDNESS:single  
820 (D) TOPOLOGY:linear  
821  
822 (ii) MOLECULE TYPE: other nucleic acid  
823 (A) DESCRIPTION: /desc = "oligonucleotide"  
824  
825 (iii) HYPOTHETICAL: YES  
826  
827 (ix) SEQUENCE DESCRIPTION:SEQ ID NO:42:  
828  
829 CCGGATAATA CGACTCACTA TAGGGCGAGC TCGGTACCCG GGTCTAGAGT CGACCTGCAG 60  
830  
831 GCATGCT 67  
832  
833  
--> 834 SEQ ID: PAGE 16  
835  
836

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**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/978,636**DATE: 11/05/98  
TIME: 14:41:34*INPUT SET: S29661.raw*

Line	Error	Original Text
38	Wrong Classification	(C)CLASSIFICATION:Not Yet Known
817	Entered (67) and Calc. Seq. Length (71) differ	(A)LENGTH:67 base pairs
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	Wrong Nucleic Acid Designator	SEQ ID: PAGE 16
834	# of Sequences for line conflicts w/ running total	SEQ ID: PAGE 16